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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/058,291

DATE: 02/10/2002
TIME: 15:09:10

#2

Input Set : N:\Crf3\RULE60\10058291.txt
Output Set: N:\CRF3\02102002\J058291.raw

SEQUENCE LISTING

4 (1) GENERAL INFORMATION:
6 (i) APPLICANT: Hartley, James L.
7 Brasch, Michael A.
9 (ii) TITLE OF INVENTION: Recombinational Cloning Using Engineered
10 Recombination Sites
12 (iii) NUMBER OF SEQUENCES: 35
14 (iv) CORRESPONDENCE ADDRESS:
15 (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C
16 (B) STREET: 1100 New York Ave., N. W. Suite 600
17 (C) CITY: Washington
18 (D) STATE: DC
19 (E) COUNTRY: USA
20 (F) ZIP: 20005-3934
22 (v) COMPUTER READABLE FORM:
23 (A) MEDIUM TYPE: Floppy disk
24 (B) COMPUTER: IBM PC compatible
25 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
26 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
28 (vi) CURRENT APPLICATION DATA:
C--> 29 (A) APPLICATION NUMBER: US/10/058,291
C--> 30 (B) FILING DATE: 30-Jan-2002
56 (C) CLASSIFICATION:
C--> 53 (vii) PRIOR APPLICATION DATA:
34 (A) APPLICATION NUMBER: 09/432,085
35 (B) FILING DATE: 1999-11-02
39 (A) APPLICATION NUMBER: 09/233,493
40 (B) FILING DATE: 20-JAN-1999
44 (A) APPLICATION NUMBER: 09/005,476
45 (B) FILING DATE: 12-JAN-1998
49 (A) APPLICATION NUMBER: 08/663,002
50 (B) FILING DATE: 07-JUN-1996
54 (A) APPLICATION NUMBER: 08/486,139
55 (B) FILING DATE: 07-JUN-1995
58 (ix) TELECOMMUNICATION INFORMATION:
59 (A) TELEPHONE: 202-371-2600
60 (B) TELEFAX: 202-371-2540
63 (2) INFORMATION FOR SEQ ID NO: 1:
65 (i) SEQUENCE CHARACTERISTICS:
66 (A) LENGTH: 25 base pairs
67 (B) TYPE: nucleic acid
68 (C) STRANDEDNESS: both
69 (D) TOPOLOGY: both

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71      (ii) MOLECULE TYPE: cDNA
76      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
78 RKYCWGCTTT YKTRTACNAA STSGB
80 (2) INFORMATION FOR SEQ ID NO: 2:
82      (i) SEQUENCE CHARACTERISTICS:
83          (A) LENGTH: 25 base pairs
84          (B) TYPE: nucleic acid
85          (C) STRANDEDNESS: both
86          (D) TOPOLOGY: both
88      (ii) MOLECULE TYPE: cDNA
93      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
95 AGCCWGCTTT YKTRTACNAA CTSGB
97 (2) INFORMATION FOR SEQ ID NO: 3:
99      (i) SEQUENCE CHARACTERISTICS:
100          (A) LENGTH: 25 base pairs
101          (B) TYPE: nucleic acid
102          (C) STRANDEDNESS: both
103          (D) TOPOLOGY: both
105      (ii) MOLECULE TYPE: cDNA
110      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
112 GTTCAGCTTT CKTRTACNAA CTSGB
114 (2) INFORMATION FOR SEQ ID NO: 4:
116      (i) SEQUENCE CHARACTERISTICS:
117          (A) LENGTH: 25 base pairs
118          (B) TYPE: nucleic acid
119          (C) STRANDEDNESS: both
120          (D) TOPOLOGY: both
122      (ii) MOLECULE TYPE: cDNA
127      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
129 AGCCWGCTTT CKTRTACNAA GTSGB
131 (2) INFORMATION FOR SEQ ID NO: 5:
133      (i) SEQUENCE CHARACTERISTICS:
134          (A) LENGTH: 25 base pairs
135          (B) TYPE: nucleic acid
136          (C) STRANDEDNESS: both
137          (D) TOPOLOGY: both
139      (ii) MOLECULE TYPE: cDNA
144      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
146 GTTCAGCTTT YKTRTACNAA GTSGB
148 (2) INFORMATION FOR SEQ ID NO: 6:
150      (i) SEQUENCE CHARACTERISTICS:
151          (A) LENGTH: 25 base pairs
152          (B) TYPE: nucleic acid
153          (C) STRANDEDNESS: both
154          (D) TOPOLOGY: both
156      (ii) MOLECULE TYPE: cDNA
161      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
163 AGCCTGCTTT TTTGTACAAA CTTGT
165 (2) INFORMATION FOR SEQ ID NO: 7:

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167      (i) SEQUENCE CHARACTERISTICS:
168          (A) LENGTH: 25 base pairs
169          (B) TYPE: nucleic acid
170          (C) STRANDEDNESS: both
171          (D) TOPOLOGY: both
173      (ii) MOLECULE TYPE: cDNA
178      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
180 AGCCTGCTTT CTTGTACAAA CTTGT
182 (2) INFORMATION FOR SEQ ID NO: 8:
184      (i) SEQUENCE CHARACTERISTICS:
185          (A) LENGTH: 25 base pairs
186          (B) TYPE: nucleic acid
187          (C) STRANDEDNESS: both
188          (D) TOPOLOGY: both
190      (ii) MOLECULE TYPE: cDNA
195      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
197 ACCCAGCTTT CTTGTACAAA CTTGT
199 (2) INFORMATION FOR SEQ ID NO: 9:
201      (i) SEQUENCE CHARACTERISTICS:
202          (A) LENGTH: 25 base pairs
203          (B) TYPE: nucleic acid
204          (C) STRANDEDNESS: both
205          (D) TOPOLOGY: both
207      (ii) MOLECULE TYPE: cDNA
212      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
214 GTTCAGCTTT TTTGTACAAA CTTGT
216 (2) INFORMATION FOR SEQ ID NO: 10:
218      (i) SEQUENCE CHARACTERISTICS:
219          (A) LENGTH: 25 base pairs
220          (B) TYPE: nucleic acid
221          (C) STRANDEDNESS: both
222          (D) TOPOLOGY: both
224      (ii) MOLECULE TYPE: cDNA
229      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
231 GTTCAGCTTT CTTGTACAAA CTTGT
233 (2) INFORMATION FOR SEQ ID NO: 11:
235      (i) SEQUENCE CHARACTERISTICS:
236          (A) LENGTH: 25 base pairs
237          (B) TYPE: nucleic acid
238          (C) STRANDEDNESS: both
239          (D) TOPOLOGY: both
241      (ii) MOLECULE TYPE: cDNA
246      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
248 GTTCAGCTTT CTTGTACAAA GTTGG
250 (2) INFORMATION FOR SEQ ID NO: 12:
252      (i) SEQUENCE CHARACTERISTICS:
253          (A) LENGTH: 25 base pairs
254          (B) TYPE: nucleic acid
255          (C) STRANDEDNESS: both

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256         (D) TOPOLOGY: both
258     (ii) MOLECULE TYPE: cDNA
263     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
265 AGCCTGCTTT TTTGTACAAA GTTGG
267 (2) INFORMATION FOR SEQ ID NO: 13:
269     (i) SEQUENCE CHARACTERISTICS:
270         (A) LENGTH: 25 base pairs
271         (B) TYPE: nucleic acid
272         (C) STRANDEDNESS: both
273         (D) TOPOLOGY: both
275     (ii) MOLECULE TYPE: cDNA
280     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
282 AGCCTGCTTT CTTGTACAAA GTTGG
284 (2) INFORMATION FOR SEQ ID NO: 14:
286     (i) SEQUENCE CHARACTERISTICS:
287         (A) LENGTH: 25 base pairs
288         (B) TYPE: nucleic acid
289         (C) STRANDEDNESS: both
290         (D) TOPOLOGY: both
292     (ii) MOLECULE TYPE: cDNA
297     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
299 ACCCAGCTTT CTTGTACAAA GTTGG
301 (2) INFORMATION FOR SEQ ID NO: 15:
303     (i) SEQUENCE CHARACTERISTICS:
304         (A) LENGTH: 25 base pairs
305         (B) TYPE: nucleic acid
306         (C) STRANDEDNESS: both
307         (D) TOPOLOGY: both
309     (ii) MOLECULE TYPE: cDNA
314     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
316 GTTCAGCTTT TTTGTACAAA GTTGG
318 (2) INFORMATION FOR SEQ ID NO: 16:
320     (i) SEQUENCE CHARACTERISTICS:
321         (A) LENGTH: 25 base pairs
322         (B) TYPE: nucleic acid
323         (C) STRANDEDNESS: both
324         (D) TOPOLOGY: both
326     (ii) MOLECULE TYPE: cDNA
331     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
333 GTTCAGCTTT CTTGTACAAA GTTGG
335 (2) INFORMATION FOR SEQ ID NO: 17:
337     (i) SEQUENCE CHARACTERISTICS:
338         (A) LENGTH: 39 base pairs
339         (B) TYPE: nucleic acid
340         (C) STRANDEDNESS: both
341         (D) TOPOLOGY: both
343     (ii) MOLECULE TYPE: cDNA
348     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
350 CCACCACAAA CGCGTCCATG GAATTACACT TTAATTTAG

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Input Set : N:\Crf3\RULE60\10058291.txt

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352 (2) INFORMATION FOR SEQ ID NO: 18:
354     (i) SEQUENCE CHARACTERISTICS:
355         (A) LENGTH: 39 base pairs
356         (B) TYPE: nucleic acid
357         (C) STRANDEDNESS: both
358         (D) TOPOLOGY: both
360     (ii) MOLECULE TYPE: cDNA
365     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
367 CCACCACAAG TCGACGCATG CCGACAGCCT TCCAAATGT      39
369 (2) INFORMATION FOR SEQ ID NO: 19:
371     (i) SEQUENCE CHARACTERISTICS:
372         (A) LENGTH: 46 base pairs
373         (B) TYPE: nucleic acid
374         (C) STRANDEDNESS: both
375         (D) TOPOLOGY: both
377     (ii) MOLECULE TYPE: cDNA
382     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
384 GGCCGATTAC GATATCCCAA CGACCGAAAA CCTGTATTTT CAGGGT      46
386 (2) INFORMATION FOR SEQ ID NO: 20:
388     (i) SEQUENCE CHARACTERISTICS:
389         (A) LENGTH: 30 base pairs
390         (B) TYPE: nucleic acid
391         (C) STRANDEDNESS: both
392         (D) TOPOLOGY: both
394     (ii) MOLECULE TYPE: cDNA
399     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
401 CAGGTTTTTCG GTCGTTGGGA TATCGTAATC      30
403 (2) INFORMATION FOR SEQ ID NO: 21:
405     (i) SEQUENCE CHARACTERISTICS:
406         (A) LENGTH: 47 base pairs
407         (B) TYPE: nucleic acid
408         (C) STRANDEDNESS: both
409         (D) TOPOLOGY: both
411     (ii) MOLECULE TYPE: cDNA
416     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
418 GGCCAGATTA CGATATCCCA ACGACCGAAA ACCTGTATTT TCAGGGT      47
420 (2) INFORMATION FOR SEQ ID NO: 22:
422     (i) SEQUENCE CHARACTERISTICS:
423         (A) LENGTH: 31 base pairs
424         (B) TYPE: nucleic acid
425         (C) STRANDEDNESS: both
426         (D) TOPOLOGY: both
428     (ii) MOLECULE TYPE: cDNA
433     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
435 CAGGTTTTTCG GTCGTTGGGA TATCGTAATC T      31
437 (2) INFORMATION FOR SEQ ID NO: 23:
439     (i) SEQUENCE CHARACTERISTICS:
440         (A) LENGTH: 48 base pairs
441         (B) TYPE: nucleic acid

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VERIFICATION SUMMARY

DATE: 02/10/2002

PATENT APPLICATION: US/10/058,291

TIME: 15:09:11

Input Set : N:\Crf3\RULE60\10058291.txt

Output Set: N:\CRF3\02102002\J058291.raw

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:33 M:220 C: Keyword misspelled or invalid format, [(vii) PRIOR APPLICATION DATA:]